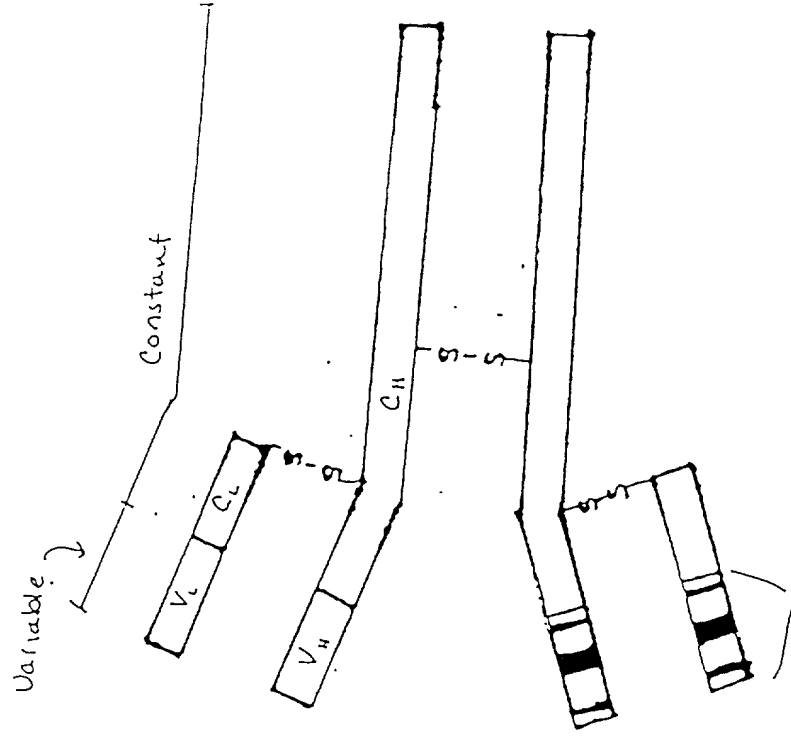


FIGURE 2



Hypervariable regions
Containing 3 CDRs

FIGURE 3

Effect of Anti-Staph MAB 96-110 on Survival in a Lethal *S. aureus* Sepsis Model

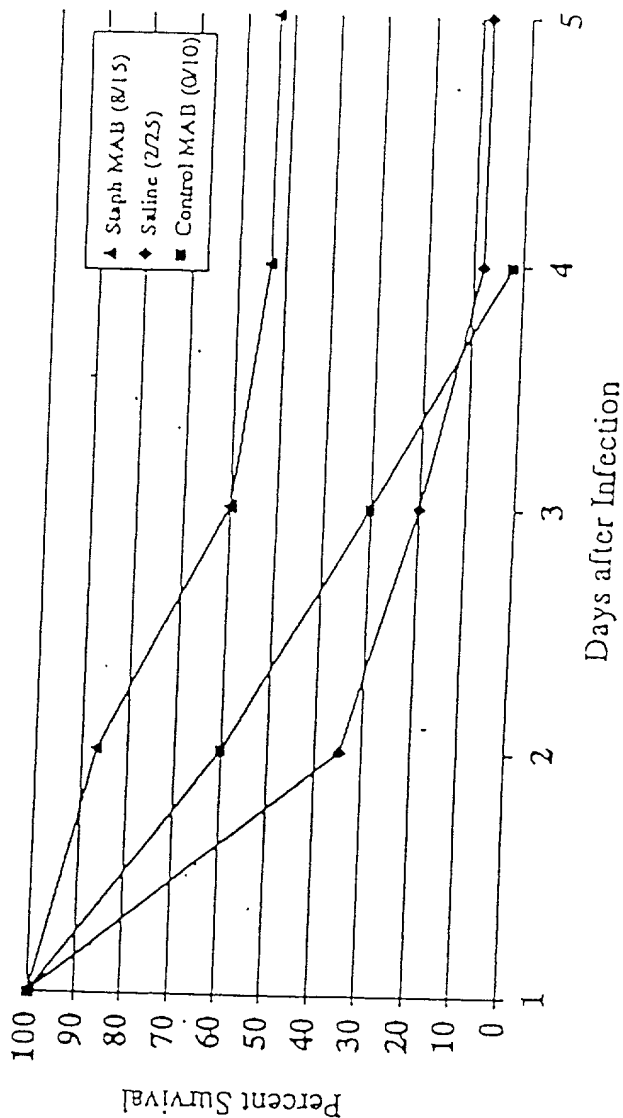


FIGURE 4

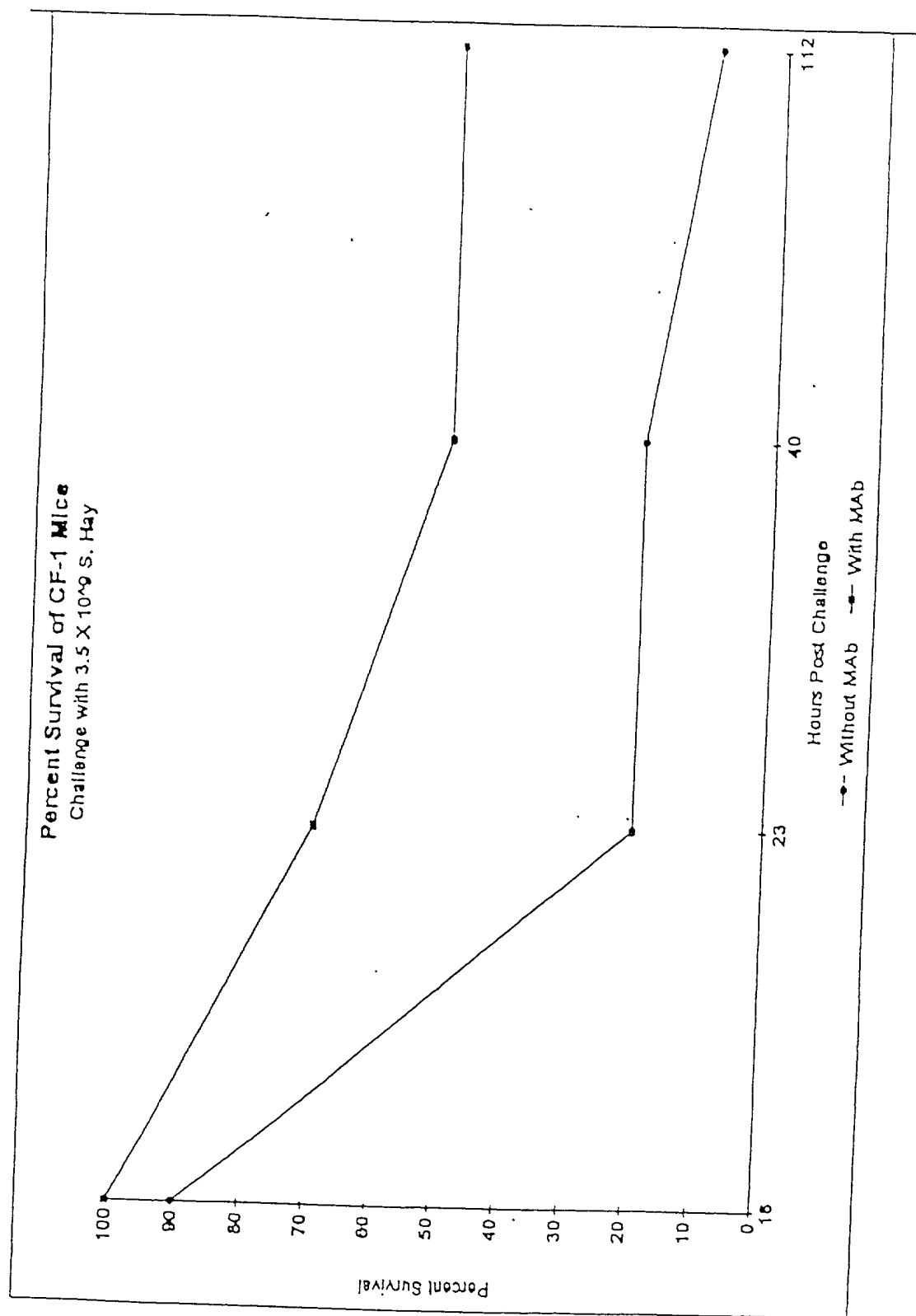


FIGURE 5

6MER.SEQ		10	20	30	
11	13.6mer2-1	GGGGCTCATC	CGGATAGGGT	TTATGGGGCC	SEQ ID NO. 4
61		C A H A D R V	Y G A		SEQ ID NO. 5
42	14.6mer2-2	GGGGCTCATC	CGGATAGGGT	TTATGGGGCC	
65		C A H A D R V	Y G A		
43	15.6mer2-3	GGGGCTCATC	CGGATAGGGT	TTATGGGGCC	
66		C A H A D R V	Y G A		
44	16.6mer2-4	GGGGCTCATC	CGGATAGGGT	TTATGGGGCC	SEQ ID NO. 6
62		C A H A D R V	Y G A		SEQ ID NO. 7
45	17.6mer2-5	GGGGCTCATC	CGGATAGGGT	TTATGGGGCC	
67		C A H A D R V	Y G A		
46	18.6mer2-6	GGGGCTCATC	CGGATAGGGT	TTATGGGGCC	
68		C A H A D R V	Y G A		
47	19.6mer2-7	GGGGCTCATC	CGGATAGGGT	TTATGGGGCC	
69		C A H A D R V	Y G A		
48	20.6mer2-8	GGGGCTCATC	CGGATAGGGT	TTATGGGGCC	
70		C A H A D R V	Y G A		
49	21.6mer2-9	GGGGCTCATC	CGGATAGGGT	TTATGGGGCC	
71		C A H A D R V	Y G A		
51	23.6mer2-1	GGGGCTCATC	CGGATAGGGT	TTATGGGGCC	
72		C A H A D R V	Y G A		
52	24.6mer2-1	GGGGCTCATC	CGGATAGGGT	TTATGGGGCC	
73		C A H A D R V	Y G A		
53	25.6mer2-1	GGGGCTCATC	CGGATAGGGT	TTATGGGGCC	
74		C A H A D R V	Y G A		
54	26.6mer2-1	GGGGCTCATC	CGGATAGGGT	TTATGGGGCC	
75		C A H A D R V	Y G A		
55	27.6mer2-1	GGGGCTCATC	CGGATAGGGT	TTATGGGGCC	
76		C A H A D R V	Y G A		
56	28.6mer2-1	GGGGCTCATC	CGGATAGGGT	TTATGGGGCC	
77		C A H A D R V	Y G A		
58	30.6mer2-1	GGGGCTCATC	CGGATAGGGT	TTATGGGGCC	
78		C A H A D R V	Y G A		
59	31.6mer2-1	GGGGCTCATC	CGGATAGGGT	TTATGGGGCC	
79		C A H A D R V	Y G A		
60	32.6mer2-2	GGGGCTCATC	CGGATAGGGT	TTATGGGGCC	
80		C A H A D R V	Y G A		

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FIGURE 6

Parameter	Unit	Value	Standard Error	95% CI	P-value
Intercept		1.00	0.00	1.00	0.00
Age	Year	0.02	0.01	-0.01, 0.05	0.15
Gender					
Male		0.05	0.02	-0.01, 0.11	0.08
Female		-0.01	0.02	-0.05, 0.03	0.85
Education	Year	0.01	0.01	-0.01, 0.03	0.45
Income	Year	0.01	0.01	-0.01, 0.03	0.45
Health					
Good		0.05	0.02	-0.01, 0.11	0.08
Fair		-0.01	0.02	-0.05, 0.03	0.85
Poor		-0.01	0.02	-0.05, 0.03	0.85
Occupation					
Manager		0.05	0.02	-0.01, 0.11	0.08
Professional		-0.01	0.02	-0.05, 0.03	0.85
Service		-0.01	0.02	-0.05, 0.03	0.85
Unemployed		-0.01	0.02	-0.05, 0.03	0.85
Retired		-0.01	0.02	-0.05, 0.03	0.85
Homemaker		-0.01	0.02	-0.05, 0.03	0.85
Student		-0.01	0.02	-0.05, 0.03	0.85
Military		-0.01	0.02	-0.05, 0.03	0.85
Other		-0.01	0.02	-0.05, 0.03	0.85
Marital Status					
Married		0.05	0.02	-0.01, 0.11	0.08
Single		-0.01	0.02	-0.05, 0.03	0.85
Divorced		-0.01	0.02	-0.05, 0.03	0.85
Widowed		-0.01	0.02	-0.05, 0.03	0.85
Never Married		-0.01	0.02	-0.05, 0.03	0.85
Religion					
Protestant		0.05	0.02	-0.01, 0.11	0.08
Catholic		-0.01	0.02	-0.05, 0.03	0.85
Jewish		-0.01	0.02	-0.05, 0.03	0.85
Muslim		-0.01	0.02	-0.05, 0.03	0.85
Other		-0.01	0.02	-0.05, 0.03	0.85
Political Affiliation					
Democrat		0.05	0.02	-0.01, 0.11	0.08
Republican		-0.01	0.02	-0.05, 0.03	0.85
Independent		-0.01	0.02	-0.05, 0.03	0.85
Other		-0.01	0.02	-0.05, 0.03	0.85
Region					
North		0.05	0.02	-0.01, 0.11	0.08
South		-0.01	0.02	-0.05, 0.03	0.85
West		-0.01	0.02	-0.05, 0.03	0.85
Midwest		-0.01	0.02	-0.05, 0.03	0.85
Other		-0.01	0.02	-0.05, 0.03	0.85
Time	Year	0.01	0.01	-0.01, 0.03	0.45
Time Squared	Year	-0.01	0.01	-0.03, 0.01	0.05
Time Cubed	Year	0.01	0.01	-0.01, 0.03	0.45
Time Squared Cubed	Year	-0.01	0.01	-0.03, 0.01	0.05
Time Squared Cubed Squared	Year	0.01	0.01	-0.01, 0.03	0.45
Time Squared Cubed Cubed	Year	-0.01	0.01	-0.03, 0.01	0.05
Time Squared Cubed Squared Cubed	Year	0.01	0.01	-0.01, 0.03	0.45
Time Squared Cubed Squared Cubed Squared	Year	-0.01	0.01	-0.03, 0.01	0.05
Time Squared Cubed Squared Cubed Squared Cubed	Year	0.01	0.01	-0.01, 0.03	0.45
Time Squared Cubed Squared Cubed Squared Cubed Squared	Year	-0.01	0.01	-0.03, 0.01	0.05
Time Squared Cubed Squared Cubed Squared Cubed Squared Cubed	Year	0.01	0.01	-0.01, 0.03	0.45
Time Squared Cubed Squared Cubed Squared Cubed Squared Cubed Squared	Year	-0.01	0.01	-0.03, 0.01	0.05
Time Squared Cubed Squared Cubed Squared Cubed Squared Cubed Squared Cubed	Year	0.01	0.01	-0.01, 0.03	0.45
Time Squared Cubed Squared Cubed Squared Cubed Squared Cubed Squared Cubed Squared	Year	-0.01	0.01	-0.03, 0.01	0.05
Time Squared Cubed Squared Cubed Squared Cubed Squared Cubed Squared Cubed Squared Cubed	Year	0.01	0.01	-0.01, 0.03	0.45
Time Squared Cubed Squared Cubed Squared Cubed Squared Cubed Squared Cubed Squared Cubed Squared	Year	-0.01	0.01	-0.03, 0.01	0.05

15MER2_SEQ		10	20	30	40	50	60	
50	07.15mer2-1	GGGGCTTGGC	ATTGGCGTCA	TCGTATTCCT	CTTCAGCTTG	CTGCTGCTCG	TGGGGCC	SEQ ID NO. 8
70		C A W H W R H R I P	L Q L A A G R	G A				SEQ ID NO. 9
52	09.15mer2-3	GGGGCTTGGC	GGCATCGTAA	TTTTTTCAT	TTTTTTCATC	GGTCTTCAT	TGGGGCC	SEQ ID NO. 10
72		G A R R H G N F S H	F F H R S L I	G A				SEQ ID NO. 11
53	10.15mer2-4	GGGGCTTGGC	GGCTTCTT	TATCTTCT	TATCTTCTC	GGGGTTCGGC	TGGGGCC	SEQ ID NO. 12
73		G A W K A L F S H	Y R P R G S A	G A				SEQ ID NO. 13
54	11.15mer2-5	GGGGCTAGGC	ATTGGCGTCA	TCGTATTCCT	CTTCAGCTTG	CTGCTGGTCG	TGGGGCC	SEQ ID NO. 14
74		G A R H W R H R I P	L Q L A A G R	G A				SEQ ID NO. 15
56	13.15mer2-7	GGGGCTTGGC	ATTGGCGTCA	TCGTATTCCT	CTTCAGCTTG	CTGCTGCTCG	TGGGGCC	SEQ ID NO. 16
76		G A W H W R H R I P	L Q L A A G R	G A				SEQ ID NO. 17
57	14.15mer2-8	GGGGCTTGGC	ATTGGCGTCA	TCGTATTCCT	CTTCAGCTTG	CTGCTGCTCG	TGGGGCC	SEQ ID NO. 18
77		G A W H W R H R I P	L Q L A A G R	G A				SEQ ID NO. 19
58	15.15mer2-9	GGGGCTCACC	TGGCTGTTT	CTATCTCTCT	TTGCTGATG	CTACTGACCT	TGGGGCC	SEQ ID NO. 20
78		G A Q V A V L Y P P	L A D A T E L	G A				SEQ ID NO. 21
59	16.15mer2-10	GGGGCTCGTC	GGCATCGTAA	TTTTTTCAT	TTTTTTCATC	GGTCTGTGAT	TGGGGCC	SEQ ID NO. 22
79		G A R R H G N F S H	F F H R S L I	G A				SEQ ID NO. 23
60	17.15mer2-11	GGGGCTCGTC	GGCATGGTAA	TTTTTTCAT	TTTTTTCATC	GGTCTGTGAT	TGGGGCC	SEQ ID NO. 24
80		G A R R H G N F S H	F F H R S L I	G A				SEQ ID NO. 25
61	18.15mer2-12	GGGGCTTGGC	GTATGTATT	TTTTCATCGT	CATCCGCATC	TTCTAGTCC	TGGGGCC	SEQ ID NO. 26
81		G A W R H Y F S H	R H A H L R S P	G A				SEQ ID NO. 27
62	19.15mer2-13	GGGGCTTGGC	GTATGTATT	TTTTCATCGT	CATCCGCATC	TTCTAGTCC	TGGGGCC	SEQ ID NO. 28
82		G A W R H Y F S H	R H A H L R S P	G A				SEQ ID NO. 29
63	20.15mer2-14	GGGGCTTGGC	GGAGTATTT	TTCTTATCAT	CATCCGCATC	TTTGTAGTCC	TGGGGCC	SEQ ID NO. 30
93		G A W R K Y F S Y H	H A H L C S P	G A				SEQ ID NO. 31
54	21.15mer2-15	GGGGCTTGGC	GTATGTATT	TTTTCATCGT	CATCCGCATC	TTCTAGTCC	TGGGGCC	SEQ ID NO. 32
94		G A W R H Y F S H	R H A H L R S P	G A				SEQ ID NO. 33
55	22.15mer2-16	GGGGCTTGGC	GTATGTATT	TTTTCATCGT	CATCCGCATC	TTCTAGTCC	TGGGGCC	SEQ ID NO. 34
95		G A W R H Y F S H	R H A H L R S P	G A				SEQ ID NO. 35
56	23.15mer2-17	GGGGCTTGGC	GTATGTATT	TTTTCATCGT	CATCCGCATC	TTCTAGTCC	TGGGGCC	SEQ ID NO. 36
96		G A W R H Y F S H	R H A H L R S P	G A				SEQ ID NO. 37
57	24.15mer2-18	GGGGCTCGTC	GGCATCGTAA	TTTTTTCAT	TTTTTTCATC	GGTCTGTGAT	TGGGGCC	SEQ ID NO. 38
97		G A R R H C N F S H	F F H R S L I	G A				SEQ ID NO. 39
58	25.15mer2-19	GGGGCTTGGC	ATTGGCGTCA	TCGTATTCCT	CTTCAGCTTG	CTGCTGGTCG	TGGGGCC	SEQ ID NO. 40
98		G A W H W R H R I P	L Q L A A G R	G A				SEQ ID NO. 41
59	26.15mer2-20	GGGGCTCGTC	GGCATCGTAA	TTTTTTCAT	TTTTTTCATC	GGTCTGTGAT	TGGGGCC	SEQ ID NO. 42
99		G A R R H G N F S H	F F H R S L I	G A				SEQ ID NO. 43

000000

5MER1.SEQ		10	20	30	40	50	60	
17	28.15mer1-2/	GGGGCTGATT	CGATTACTTT	TCATCGTCGT	CATCATGATC	GTGTTCTTTC	TGGGGCC	SEQ ID NO. 44
17		C A D	(W) I T F	(H) R R	H (H) D	(R) V L S	G A	SEQ ID NO. 45
18	29.15mer1-3/	GGGGCTGTTT	GGATTACTTT	TCATCGTCGT	CATCATGATC	GTGTTCTTTC	TGGGGCC	SEQ ID NO. 46
18		G A C	W I T F	H R R	H H D	R V L S	G A	SEQ ID NO. 47
19	32.15mer1-6/	GGGGCTGTTT	GGATTACTTT	TCATCGTCGT	CATCATGATC	GTGTTCTTTC	TGGGGCC	
19		G A G	W I T F	H R R	H H D	R V L S	G A	
20	33.15mer1-7/	GGGGCTGTTT	GGATTACTTT	TCATCGTCGT	CATCATGATC	GTGTTCTTTC	TGGGGCC	
20		G A G	W I T F	H R R	H H D	R V L S	G A	
21	34.15mer1-8/	GGGGCTGTTT	GGATTACTTT	TCATCGTCGT	CATCATGATC	GTGTTCTTTC	TGGGGCC	
21		G A G	W I T F	H R R	H H D	R V L S	G A	
22	35.15mer1-9/	GGGGCTGTTT	GGATTACTTT	TCATCGTCGT	CATCATGATC	GTGTTCTTTC	TGGGGCC	
22		G A G	W I T F	H R R	H H D	R V L S	G A	
23	36.15mer1-10/	GGGGCTGTTT	GGATTACTTT	TCATCGTCGT	CATCATGATC	GTGTTCTTTC	TGGGGCC	
23		G A G	W I T F	H R R	H H D	R V L S	G A	
24	37.15mer1-11/	GGGGCTGTTT	GGATTACTTT	TCATCGTCGT	CATCATGATC	GTGTTCTTTC	TGGGGCC	
24		G A C	W I T F	H R R	H H D	R V L S	G A	
25	38.15mer1-12/	GGGGCTGTTT	GGATTACTTT	TCATCGTCGT	CATCATGATC	GTGTTCTTTC	TGGGGCC	
25		G A C	W I T F	H R R	H H D	R V L S	G A	
26	39.15mer1-13/	GGGGCTGTTT	GGATTACTTT	TCATCGTCGT	CATCATGATC	GTGTTCTTTC	TGGGGCC	
26		C A C	W I T F	H R R	H H D	R V L S	G A	
27	20.15mer1-14/	GGGGCTGTTT	GGATTACTTT	TCATCGTCGT	CATCATGATC	GTGTTCTTTC	TGGGGCC	
27		G A G	W I T F	H R R	H H D	R V L S	G A	
28	21.15mer1-15/	GGGGCTGTTT	GGATTACTTT	TCATCGTCGT	CATCATGATC	GTGTTCTTTC	TGGGGCC	
28		G A G	W I T F	H R R	H H D	R V L S	G A	
29	22.15mer1-16/	GGGGCTGGGA	AGGCTATGTT	TAGTCATTCT	TATCGTCATC	GGGGTTCGGC	TGGGGCC	SEQ ID NO. 48
29		G A G	(K) A M F	(S) H S	Y R (H) (R) G S A	G A	SEQ ID NO. 49	
30	23.15mer1-17/	GGGGCTGTTT	GGATTACTTT	TCATCGTCGT	CATCATGATC	GTGTTCTTTC	TGGGGCC	
30		G A C	W I T F	H R R	H H D	R V L S	G A	
31	24.15mer1-18/	GGGGCTGTTT	GGATTACTTT	TCATCGTCGT	CATCATGATC	GTGTTCTTTC	TGGGGCC	
31		G A C	W I T F	H R R	H H D	R V L S	G A	
32	25.15mer1-19/	GGGGCTGTTT	GGATTACTTT	TCATCGTCGT	CATCATGATC	GTGTTCTTTC	TGGGGCC	
32		G A G	W I T F	H R R	H H D	R V L S	G A	
33	26.15mer1-20/	GGGGCTGTTT	GGATTACTTT	TCATCGTCGT	CATCATGATC	GTGTTCTTTC	TGGGGCC	
33		G A G	W I T F	H R R	H H D	R V L S	G A	

FIGURE 9

Comparison of Signals at 6.25×10^{11} vir/mL.

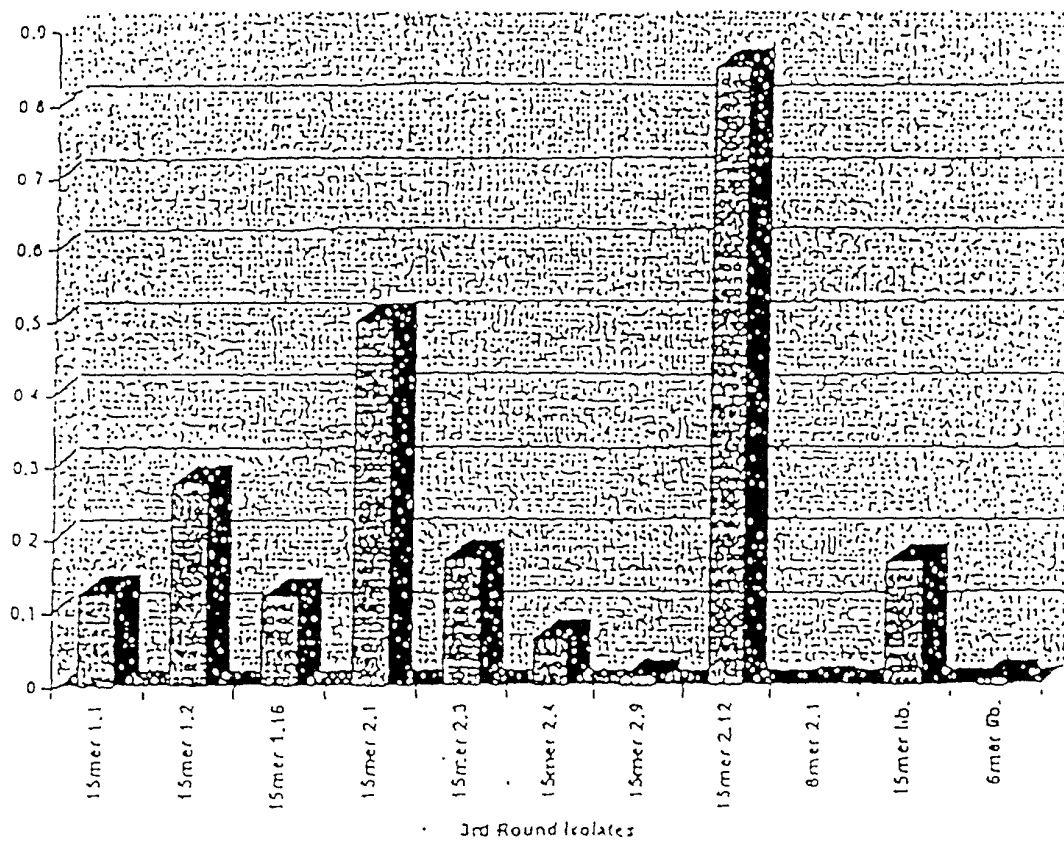


FIGURE 11

Mouse Heavy Chain "front" primers

JSS1
5'-ATTTCAGGCCCAGCCGGCCATGGCCGARGTRMAGCTKSAKGAGWC-3' SEQ ID NO 68
JSS2
5'-ATTTCAGGCCCAGCCGGCCATGGCCGARGTYCARCTKCARCARYC-3' SEQ ID NO 69
JSS3
5'-ATTTCAGGCCCAGCCGGCCATGGCCGAGGTGAAGCTKSTSGARTC-3' SEQ ID NO 70
JSS4
5'-ATTTCAGGCCCAGCCGGCCATGGCCGAVGTGMWGCTKGTGGAGWC-3' SEQ ID NO 71
JSS8
5'-ATTTCAGGCCCAGCCGGCCATGGCCGAGGTBCARCTKMARSARTC-3' SEQ ID NO 72

Mouse Heavy chain "back" primers

JS160
5'-GCTGCCACCGCCACCTGMRGAGACDGTGASTGARG-3' SEQ ID NO 73
JS161
5'-GCTGCCACCGCCACCTGMRGAGACDGTGASMTRG-3' SEQ ID NO 74
JS162
5'-GCTGCCACCGCCACCTGMRGAGACDGTGASCARG-3' SEQ ID NO 75

Mouse Light Chain Leader "front" primers

PMC12
5'-CCCCGGGCCACCATGGAGACAGACACACTCCTG-3' SEQ ID NO 76
PMC13
5'-CCCCGGGCCACCATGGATTTTCAAGTGCAGATTTTC-3' SEQ ID NO 77
PMC14
5'-CCCCGGGCCACCATGGAGWCACARWCTCAGGTC-3' SEQ ID NO 78
PMC15
5'-CCCCGGGCCACCATGKCCCCWRCTCAGYTTCTKG-3' SEQ ID NO 79
PMC55
5'-CCCCGGGCACCATGAAGTTGCCTGTTAGGCTG-3' SEQ ID NO 80

Mouse Light Chain "back" primer

OXA57
5'-GCACCTCCAGATGTTAACTGCTC-3' SEQ ID NO 81

"96-110" Specific Primers

96110HF2
5'-TAATATTCGGACAGCTACAGGTGTCCACTCCCGAAGTGATGCTGGTGGAGWCTG-3' SEQ ID NO 82
96110HB
5'-TTATAQAATTCTGAGGAGACGGTGAGTGAG-3' SEQ ID NO 83
96110BLF
5'-TTAGGCQAATATCGTTCTCTCCAGTCTCC-3' SEQ ID NO 84
96110BLB
5'-GTAACCGTTCGAAAAGTGTACTTACGTTTTATTTCAGCATGGTCC-3' SEQ ID NO 85

099315-062901

FIGURE 12

96-110 anti-staph (HAY) heavy chain variable region (type IIIA)

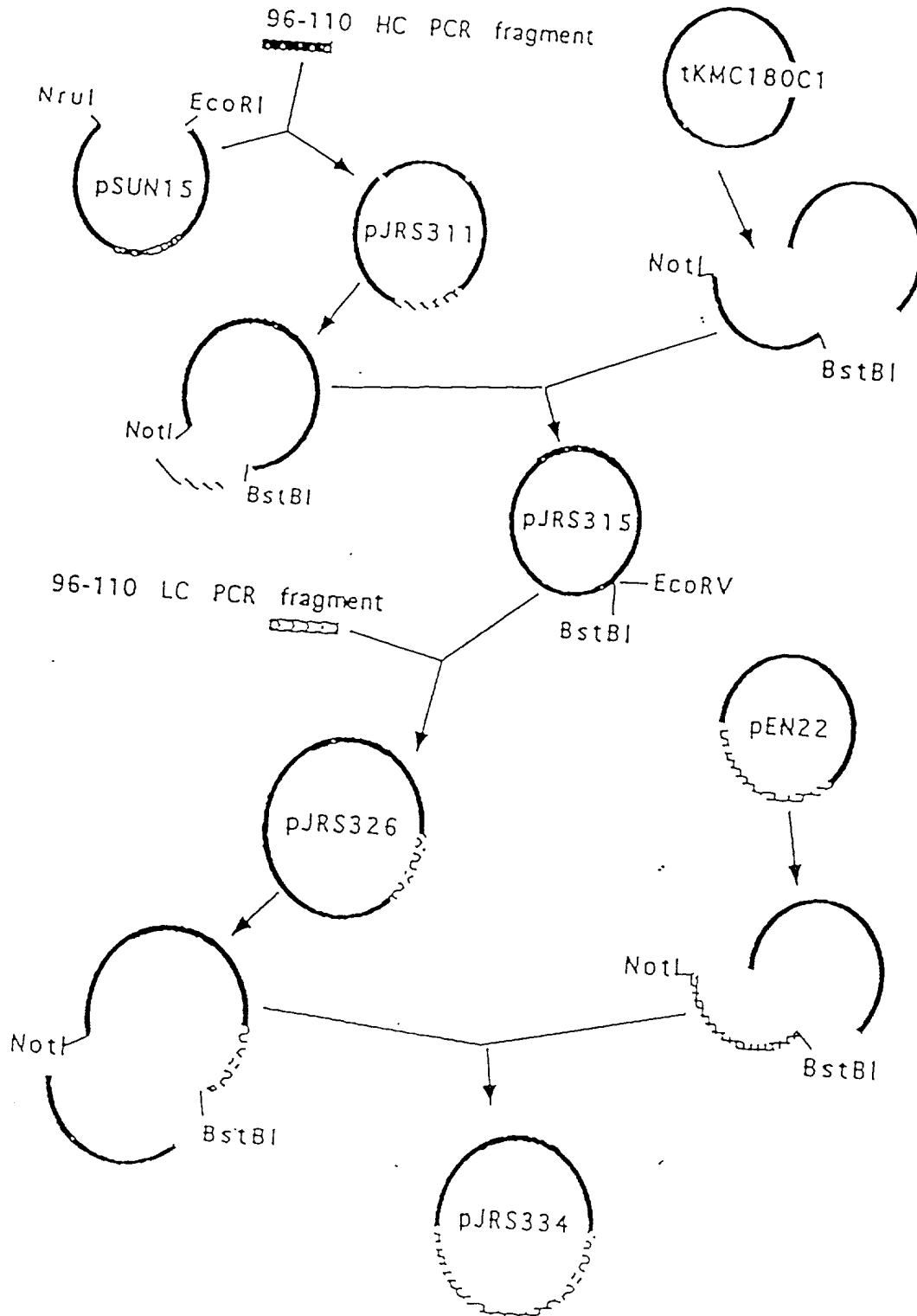
GAAGTCATGCTGGTGGAGTCTGGTGGAGATGGTGGCAGCCTAAAGGGTCATTTGAACTCTCATGTGCAAGCTCTGCACTTCATCAT 3EQ 1D NO. 86
 E V M L V E S G G G L V Q P K G S L K L S C A A S G F T F H SEQ 1D NO. 87
 AACTAGCCCATGAAAT TGGGTCCGCGCAGGCTCCAGGAAGGGTTGGAAATGGTTGCT SEQ 1D NO. 88
 N Y A H N W V R Q A P G K G L E H V A SEQ 1D NO. 89
 CCGATAGAGAACTAAAGTAATTAATTCGCAACATTTTAAAGCCGATTCAGTGAAGAC SEQ 1D NO. 90
 R I R S K S N N Y A T P Y A D S V X D SEQ 1D NO. 91
 AGGTTCAACCATCTCCAGACATGATTCACAAAGCAATGCTCTATCTGCAATTCGAAACATTCGAAACTGAGGACACAGCCATGTATTACTGTGTACAG SEQ 1D NO. 92
 R P T I S R D D S Q S H L Y L Q H N N L K T E D T A H Y Y C V R SEQ 1D NO. 93
 CCGGCGGCTTCAGGGATTGACTATGCTATGGACTAC TGGGTTCAGGAAGCCTCACTACCGTCTCTCA SEQ 1D NO. 94
 R G A S G I D Y A H D Y H G Q G T S L T V S S SEQ 1D NO. 95

96-110 anti-staph (HAY) light chain variable region (type VI)

CAAAATGTTCTCTCCAGTCTCCAGCAATCTCTGTCTGCACTCTCCAGGGGAAAGGTCAATGACTTCC SEQ 1D NO. 96
 Q I V L S Q S P A I L S A S P G E K V T H T C SEQ 1D NO. 97
 ACGGCGCAGCTCAAGCTTAAATTAATCCAG SEQ 1D NO. 98
 R A S S S V N Y H R SEQ 1D NO. 99
 TGGTACCAGCAGAGCCAGGATCTCTCCGCAACCCCTGGATTCT GCGACATCCAACTGGTTCT SEQ 1D NO. 100
 H Y Q Q Q K P G S S P K P H I S A T S N L A S SEQ 1D NO. 101
 CGAGTCCCTGCTCGCTTCAGTGGCAGTGGGTCTGGGACCTCTTACTCTCTCAATCAAGAGTGGAGCCTGAAATGCTGCACTTATTACTCC SEQ 1D NO. 102
 G V P A R P S G S G S G T S Y S L T I S R V E A E D A A T Y Y C SEQ 1D NO. 103
 CAGCAGTGGAGTAGTAAGCCAGCCAGC TCCGAGGGGGGACCATGCTGGAATAGA SEQ 1D NO. 104
 Q Q W S S N P P T F G G G T H L E I R SEQ 1D NO. 105

CDR Regions Underlined

FIGURE 13



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FIGURE 14

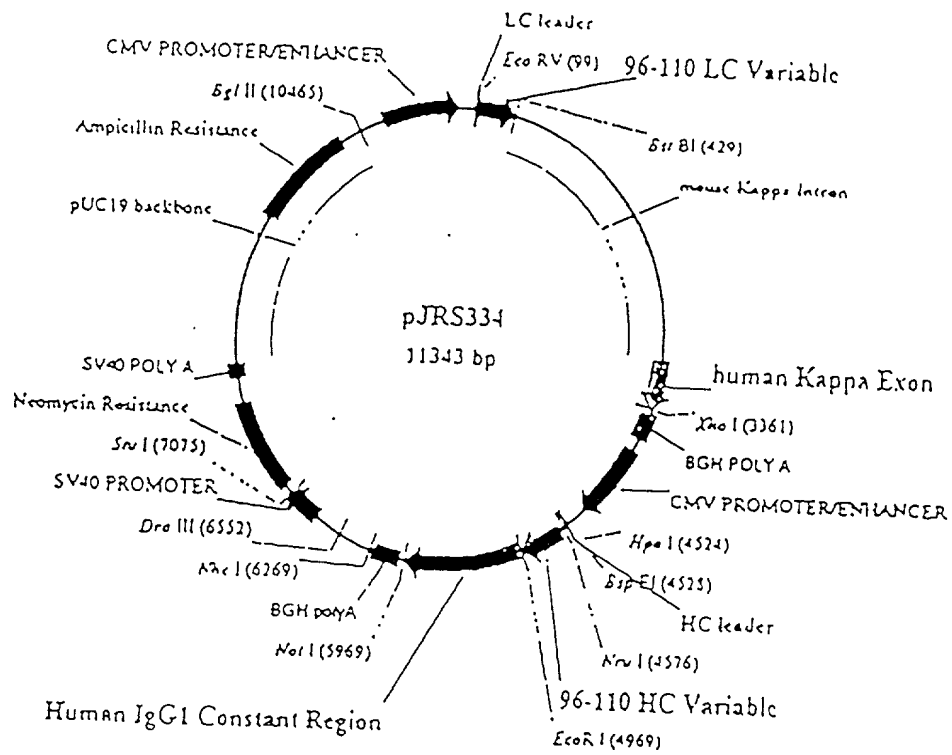


FIGURE 15: Antibody Production ELISA

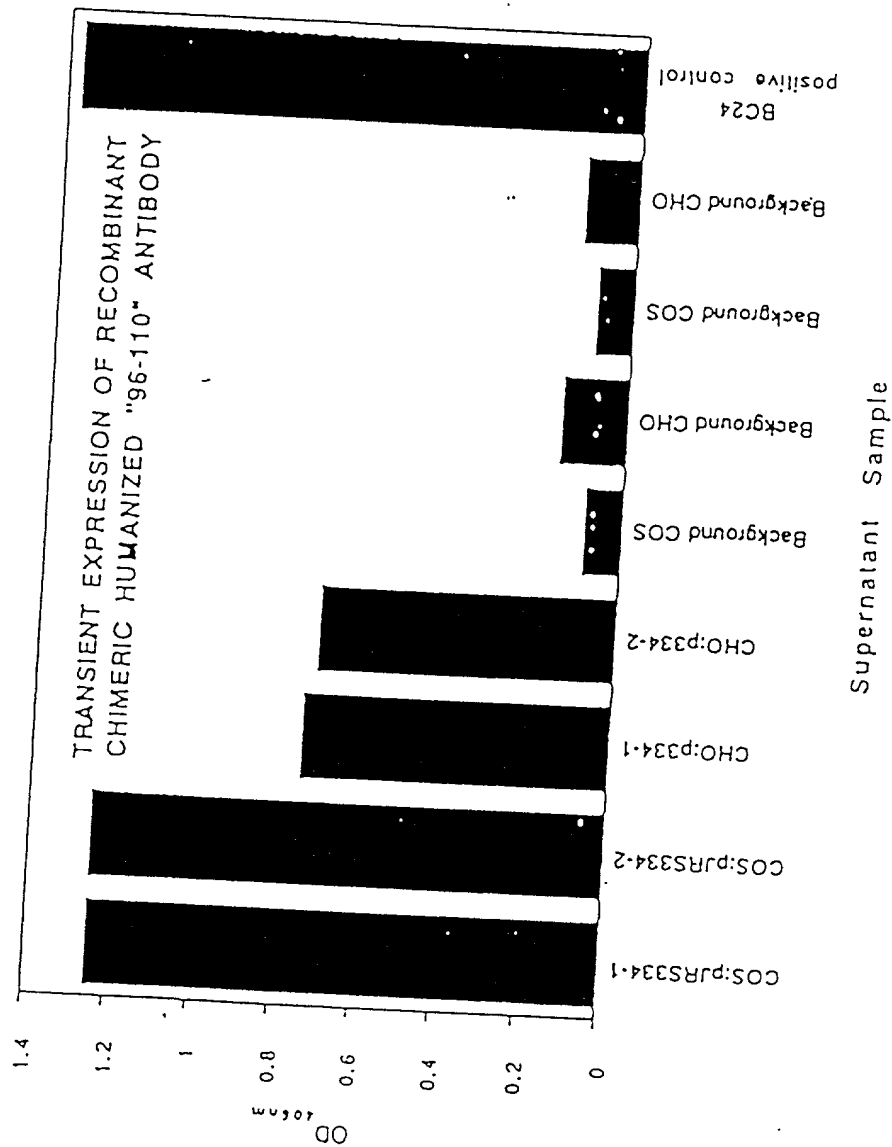
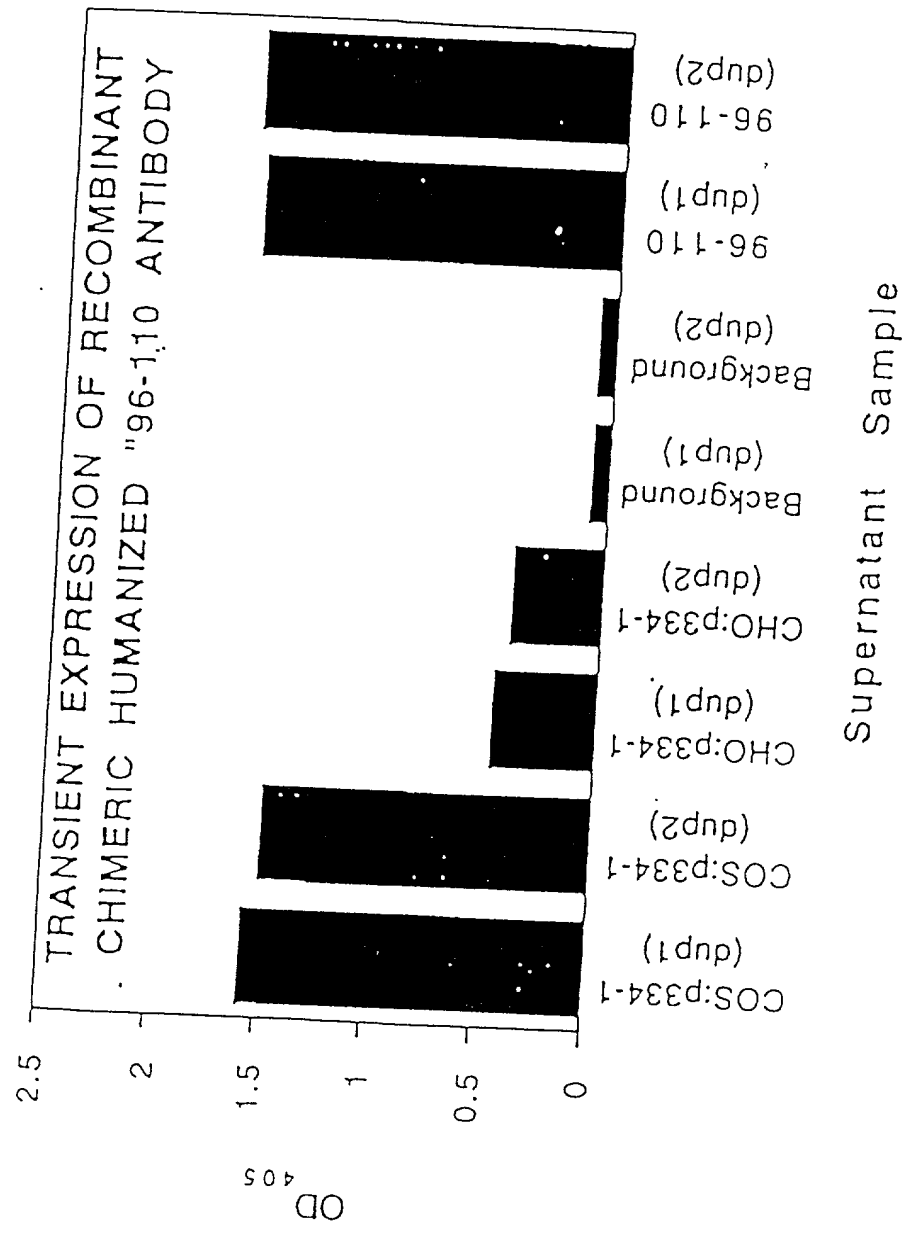
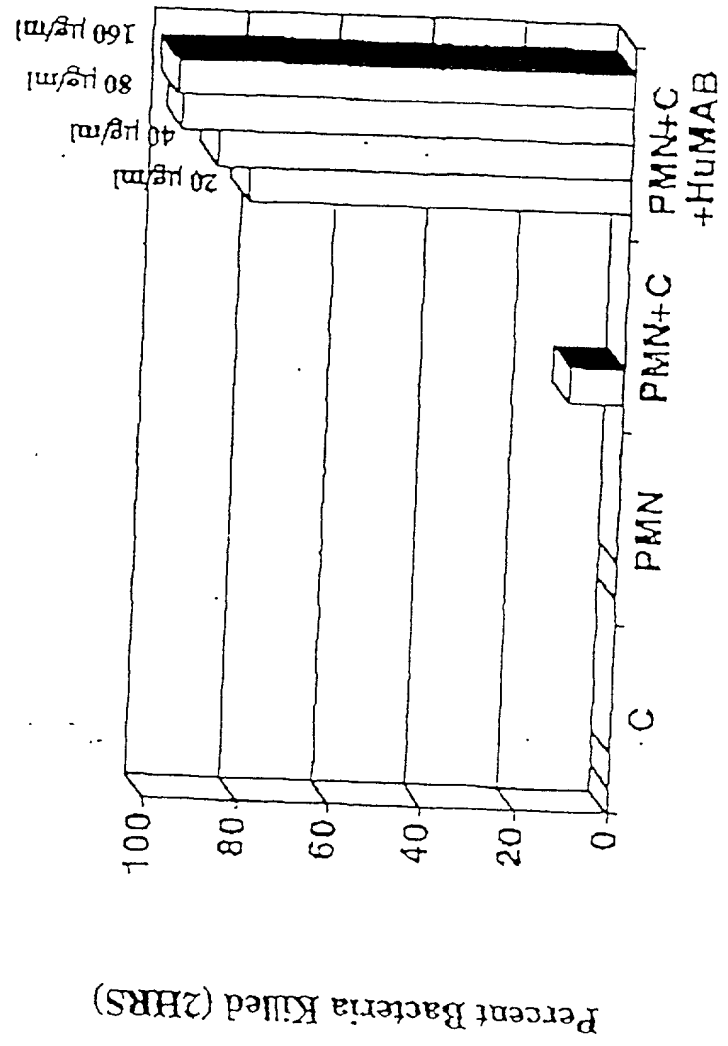


FIGURE 16: Anti-Staph HAY Activity ELISA

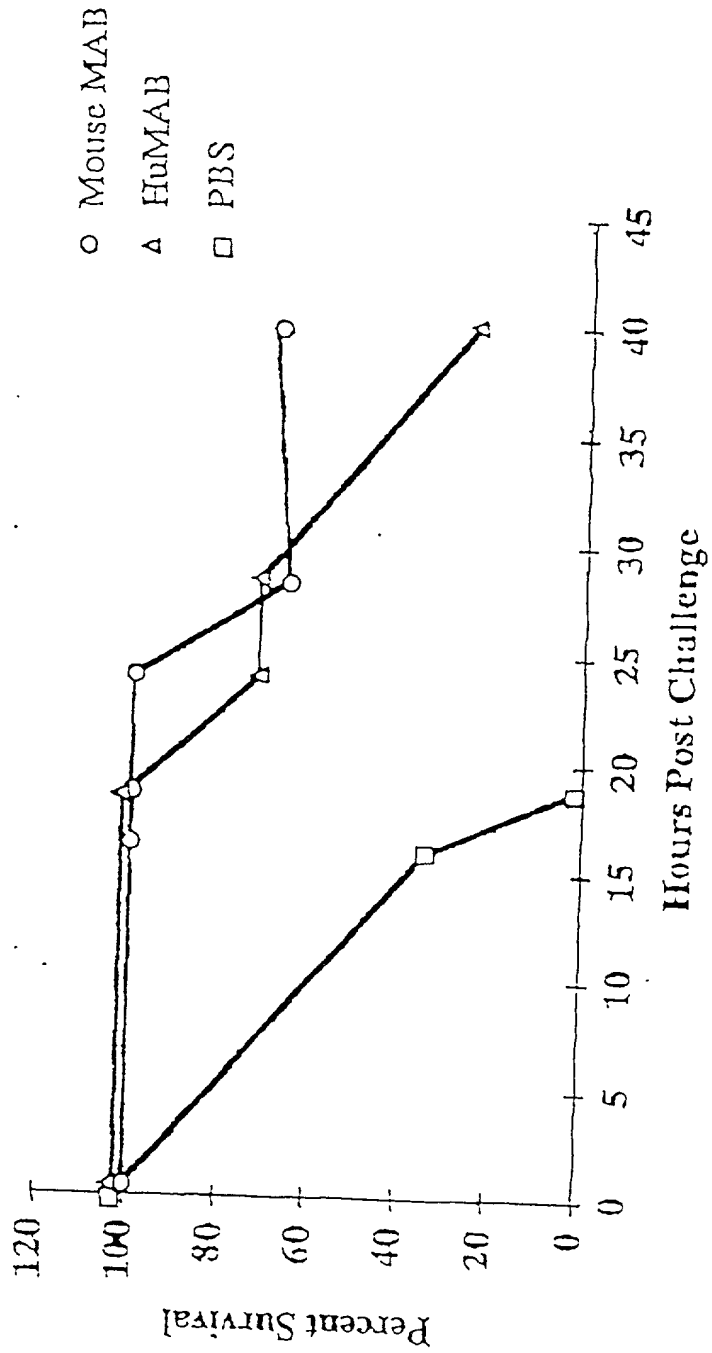


Opsonic Activity of HuMAB 96-110 for *S.epidermidis* in a Neutrophil Mediated Opsonophagocytic Bactericidal Assay Using Human Complement



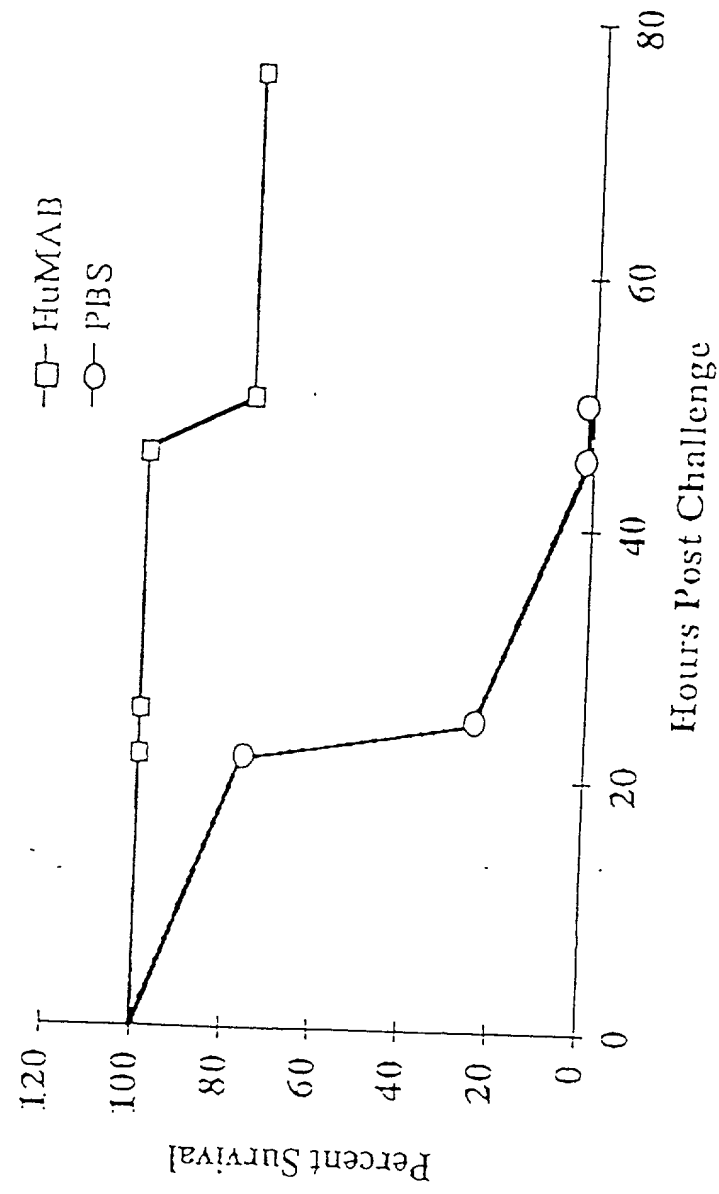
C-Barb-Ex (1:4), Human
PMN-Human
Bacteria-*S.epidermidis* (Strain Hay)

Pilot Study to Compare the Effect of Mouse MAB 96-110 and HuMAB 96-110 in a Lethal Model of *S. Epidermidis* Sepsis



MAB dose: 14 mg/kg given IP, 24 and 1 Hour prior to infection

Survival of CF-1 Mice after Intraperitoneal Challenge with 3×10^9 *S. epidermidis* (Hay)



18 mg/kg/dose, IP, 24 and 1 Hour prior to infection

Effect of HuMAB 96-110 on Bacteremia in a Lethal *S. epidermidis* Sepsis Model

Geometric Mean Bacteremia Level

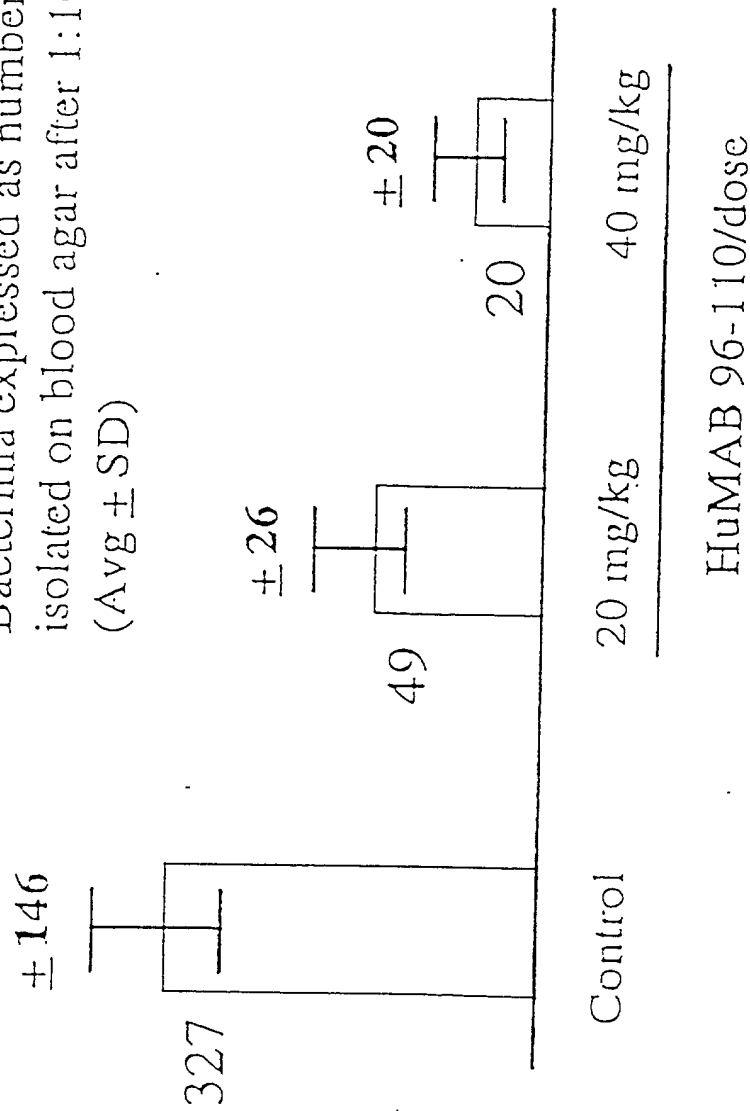
Saline				
Placebo	6.5×10^4	7.2×10^4	5.2×10^4	7×10^3
HuMAB				
96-110	3×10^2	7.5×10^2	2.1×10^1	1.7×10^1
	4 hrs	8 hrs	12 hrs	18 hrs

Time Post Infection

HuMAB 96-110 18 mg/kg/dose or saline given IP, 24 and 1 hour
prior to IP infection with 3×10^9 *S. epidermidis* (Hay)

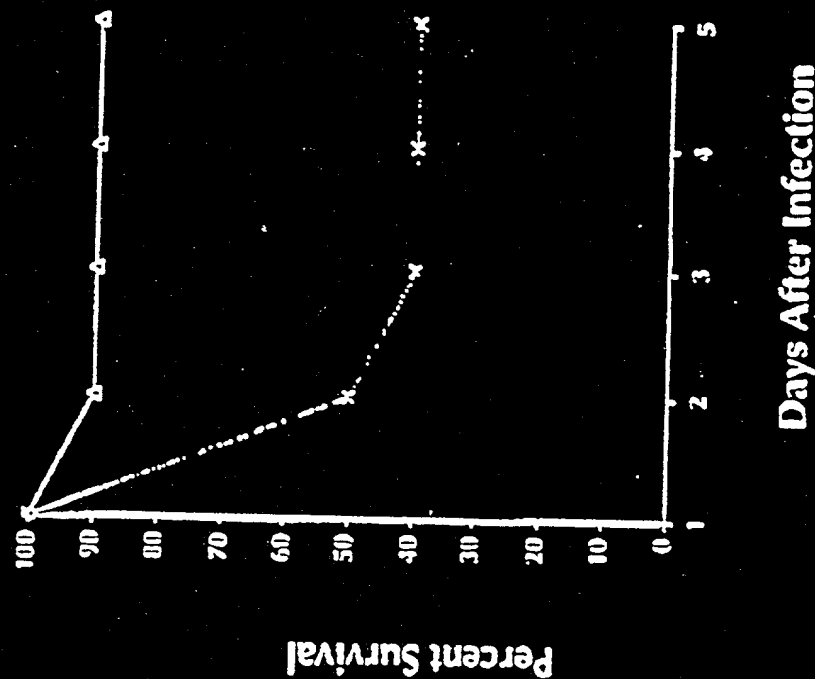
Bacteremia levels 4 hrs after infection with 3×10^9 *S.epidermidis**

Bactermia expressed as number of bacteria
isolated on blood agar after 1:1000 dilution
(Avg \pm SD)



* CF-1 mice infected IP with strain Hay-HuMAB given IP x 2

The Effect of Hu 96-110 on Survival in a Lethal Neonatal *S.epidermidis** Sepsis Model: Study II



Survival:
Hu 96-110 27/30 (90%)
Saline 12/30 (40%)

—○— Hu 96-110
...x... Saline Control
* Different SE Strains

Study II:
Infection - ~5x10⁷ SE, SQ (with plastic catheter SQ)
Lipid Emulsion - 0.2 ml, 20% IP day -1 and +1, 2
doses day 0
HuMAB or Saline- 0.2 ml, IP 30 min before and 24 hrs
after infection
Dose: 50-60 mg/kg per dose